```
SEQUENCE LISTING
<110 >Tonen Corporation
<120 >Method for Detection or Measurement of Hepatitis C V
<160 >8
<210 >1
<211 >177
<212 >PRT
<213 >Hepatitis C virus
<400 >1
  Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Pro Glu
                                                        15
                                     10
  Phe Met Gly Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr
              20
                                 25
  Asn Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val
                             40
  Gly Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg
      50
                          55
  Ala Thr Arg Lys Thr Ser Lys Arg Ser Gln Pro Arg Gly Gly Arg Arg
                                         75
  Pro Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro
                                     90
                  85
  Gly Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly
                                105
             100
  Trp Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp
                                                125
                             120
  Pro Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr
      130
                         135
  Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe
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155

175

Leu Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu

170

150

165

Asp

```
<211 >160
<212 >TRP
<213 >Hepatitis C virus
<400 >2
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                                     10
 Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
                                 25
              20
 Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
                             40
 Thr Arg Lys Thr Ser Lys Arg Ser Gln Pro Arg Gly Gly Arg Arg Pro
    . 50
                          55
 Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
                      70
                                         75
  Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
                                                         95
                                     90
                  85
  Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
                                                    110
             1.00
                                 105
  Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
                             120
                                                125
  Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe Leu
                                           140
                       135
    130
  Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
                                                            160
                                         155
                     150
  145
<210 >3
<211 >20
<212 >PRT
<213 >Artificial Sequence
<220 >
<223 >
<400 >3
  Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly Gly Val Tyr Leu
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10

5

15

<210 >2

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Leu Pro Arg Arg
              20
<210 >4
<211 >10
<212 >PRT
<213 >Artificial Sequence
<220 >
<223 >
<400 >4
 Gly Pro Arg Leu Gly Val Arg Ala Thr Arg
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<210 >5
<211 >21
<212 >PRT
<213 >Artificial Sequence
<220 >
<223 >
<400 >5
 Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro Arg His Arg
    1
                   5
                                     10
                                                        15
  Ser Arg Asn Val Gly
              20
<210 >6
<211 >20
<212 >PRT
<213 >Artificial Sequence
<220 >
<230 >
<400 >6
  Asp Pro Arg His Arg Ser Arg Asn Val Gly Lys Val Lle Asp Thr Leu
                                     10
                                                         15
    1
  Thr Cys Gly Phe
              20
<210 >7
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<211	>24
<212	>DNA
<213	>Artificial Sequence
<220	>Probe
<230	>Synthetic DNA
<400	>7
gaa	ttcatgg gcacgaatcc taaa
<210	>8
<211	>21
<212	>DNA
<213	>Artificial Sequence
<220	>Probe
<230	>Synthetic DNA
<400	>8

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